



SEQUENCE LISTING

<110> HYBRID BIOSCIENCES

Pierre, LEGRAIN

<120> Identification of the Anti-s28 Factor in Helicobacter pylori, in Campylobacter jejuni and in Pseudomonas aeruginosa and Application Thereof

<130> B4797A

<140> US 10/066,127

<141> 2002-01-31

<150> US 60/265,465

<151> 2001-01-31

<160> 29

<170> PatentIn version 3.1

<210> 1

<211> 90

<212> DNA

<213> Helicobacter pylori

<220>

<221> SID1122

<222> (1)..(90)

<223> the Selected Interacting Domain (SID®) of HP1122

<400> 1

atcaagaaag cgattgaaaa taaccagtat aaaatcaact tgcattgagac ttctcacaaa 60

atggcaaagg atttattggg gataagctag

90

<210> 2

<211> 29

<212> PRT

<213> Helicobacter pylori

<220>

<221> SID1122

<222> (1) .. (29)

<223>

<400> 2

Ile Lys Lys Ala Ile Glu Asn Asn Gln Tyr Lys Ile Asn Leu His Glu
1 5 10 15

Thr Ser His Lys Met Ala Lys Asp Leu Leu Gly Ile Ser
20 25

<210> 3

<211> 177

<212> DNA

<213> Helicobacter pylori

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<221> SID1032

<222> (1) .. (177)

<223> the Selected Interacting Domain (SID®) HP1032

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aaagcgctga atcaaattgag cgaaagagag caaatcctta tccagcttta ttactttgaa 60
gagttgaatt tgagcgagat taaagagatt ttaggcatta ctgaatcgcg cattttctcaa 120
atcattaaag aagtgattaa aaaggtgcgt aaatccttag gagtggatca tggctga 177

<210> 4

<211> 58

<212> PRT

<213> Helicobacter pylori

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<221> SID1032

<222> (1)..(58)

<223>

<400> 4

Lys Ala Leu Asn Gln Met Ser Glu Arg Glu Gln Ile Leu Ile Gln Leu
1 5 10 15

Tyr Tyr Phe Glu Glu Leu Asn Leu Ser Glu Ile Lys Glu Ile Leu Gly
20 25 30

Ile Thr Glu Ser Arg Ile Ser Gln Ile Ile Lys Glu Val Ile Lys Lys
35 40 45

Val Arg Lys Ser Leu Gly Val Asp His Gly
50 55

<210> 5

<211> 231

<212> DNA

<213> Helicobacter pylori

<220>

<221> HP1122

<222> (1)..(231)

<223> the ORF of sigma28 factor

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cagtctttgg ggaattataa gcgtgtggaa aagaatgaaa aagttgaaaa caatgaggcc 120
 gctcttgata gggtagctga gatcaagaaa gcgattgaaa ataaccagta taaaatcaac 180
 ttgcatgaga cttctcaciaa aatggcaaag gatttattgg ggataagcta g 231

<210> 6

<211> 76

<212> PRT

<213> Helicobacter pylori

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<221> HP1122

<222> (1)..(76)

<223>

<400> 6

Met Asn Ile Lys Leu Lys Asp Phe Thr Met Ile Asn Ala Val Ser Ser
 1 5 10 15

Leu Ala Pro Val Gln Ser Leu Gly Asn Tyr Lys Arg Val Glu Lys Asn
 20 25 30

Glu Lys Val Glu Asn Asn Glu Ala Ala Leu Asp Arg Val Ala Glu Ile
 35 40 45

Lys Lys Ala Ile Glu Asn Asn Gln Tyr Lys Ile Asn Leu His Glu Thr
 50 55 60

Ser His Lys Met Ala Lys Asp Leu Leu Gly Ile Ser
 65 70 75

<210> 7

<211> 768

<212> DNA

<213> Helicobacter pylori

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<221> HP1032

<222> (1)..(768)

<223> The ORF of anti sigma 28 factor

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aaaaatatag aaaaggtttt gaacgcctat gataagcaac aacaccacca tcaagacgat      120
ctcgctattc agtattttacc agccgtgcgc gccatggcgt ttcgtctaaa agagcgcttg      180
cccagctcta ttgatttttaa cgatctgggt tctattggca ctgaagaatt gattaaatta      240
gccaggcggt atgagagcgc gttaaacgat tctttttggg ggtatgcgaa gactcgtgtc      300
aatggggcgca tgtagatta tttgcgctct ttagatgtga tttctcgctc tagcaggaaa      360
ctcattaaaa gcattgatat tgaaatcacc aaacacctta atgagcatgg gaaagagcct      420
agcgatgcgt atttagcgca aacttttaggc gaaaatattg aaaaaattaa agaagccaaa      480
acggcttcag atatttatgc gttagtgcc aatgatgaac aattcaatgc gattgagcaa      540
gatgaaatca ctaaaaaaat tgaagcagaa gagttgtag agcatgtcca aaaagcgctg      600
aatcaaatga gcgaaagaga gcaaatcctt atccagcttt attactttga agagttgaat      660
ttgagcgaga ttaaagagat tttaggcatt actgaatcgc gcattttctca aatcattaaa      720
gaagtgatta aaaaggtgcg taaatcctta ggagtggatc atggctga      768
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<210> 8

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<213> Helicobacter pylori

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<221> HP1032

<222> (1)..(255)

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<400> 8

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Glu Thr Ser Glu Lys Asn Ile Glu Lys Val Leu Asn Ala Tyr Asp Lys
 20 25 30

Gln Gln His His Gln Asp Asp Leu Ala Ile Gln Tyr Leu Pro Ala
 35 40 45

Val Arg Ala Met Ala Phe Arg Leu Lys Glu Arg Leu Pro Ser Ser Ile
 50 55 60

Asp Phe Asn Asp Leu Val Ser Ile Gly Thr Glu Glu Leu Ile Lys Leu
 65 70 75 80

Ala Arg Arg Tyr Glu Ser Ala Leu Asn Asp Ser Phe Trp Gly Tyr Ala
 85 90 95

Lys Thr Arg Val Asn Gly Ala Met Leu Asp Tyr Leu Arg Ser Leu Asp
 100 105 110

Val Ile Ser Arg Ser Ser Arg Lys Leu Ile Lys Ser Ile Asp Ile Glu
 115 120 125

Ile Thr Lys His Leu Asn Glu His Gly Lys Glu Pro Ser Asp Ala Tyr
 130 135 140

Leu Ala Gln Thr Leu Gly Glu Asn Ile Glu Lys Ile Lys Glu Ala Lys
 145 150 155 160

Thr Ala Ser Asp Ile Tyr Ala Leu Val Pro Ile Asp Glu Gln Phe Asn
 165 170 175

Ala Ile Glu Gln Asp Glu Ile Thr Lys Lys Ile Glu Ala Glu Glu Leu
 180 185 190

Leu Glu His Val Gln Lys Ala Leu Asn Gln Met Ser Glu Arg Glu Gln
 195 200 205

Ile Leu Ile Gln Leu Tyr Tyr Phe Glu Glu Leu Asn Leu Ser Glu Ile
 210 215 220

Lys Glu Ile Leu Gly Ile Thr Glu Ser Arg Ile Ser Gln Ile Ile Lys
 225 230 235 240

Glu Val Ile Lys Lys Val Arg Lys Ser Leu Gly Val Asp His Gly
 245 250 255

<210> 9
 <211> 65
 <212> PRT
 <213> Campylobacter jejuni

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 <221> Cj1464
 <222> (1)..(65)
 <223> Cj1464 protein

<400> 9
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 1 5 10 15
 Thr Asn Arg Ile Asp Lys Glu Thr Lys Thr Asn Asp Thr Gln Lys Thr
 20 25 30
 Glu Asn Asp Lys Ala Ser Lys Ile Ala Glu Gln Ile Lys Asn Gly Thr
 35 40 45
 Tyr Lys Ile Asp Thr Lys Ala Thr Ala Ala Ala Ile Ala Asp Ser Leu
 50 55 60
 Ile
 65

<210> 10
 <211> 107
 <212> PRT
 <213> Pseudomonas aeruginosa

<220>
 <221> PA3351
 <222> (1)..(107)
 <223> PA3351 protein

<400> 10

Met Val Ile Asp Phe Asn Arg Leu Asn Pro Gly Ser Thr Pro Ala Thr
1 5 10 15

Thr Gly Arg Thr Gly Ser Thr Ala Ala Gly Arg Pro Asp Ala Thr Gly
20 25 30

Ala Asp Lys Ala Gly Gln Ala Ala Thr Ser Ala Pro Lys Ser Gly Glu
35 40 45

Ser Val Gln Ile Ser Glu Thr Ala Gln Asn Met Gln Lys Val Thr Asp
50 55 60

Gln Leu Gln Thr Leu Pro Val Val Asp Asn Asp Lys Val Ala Arg Ile
65 70 75 80

Lys Gln Ala Ile Ala Asp Gly Thr Tyr Gln Val Asp Ser Glu Arg Val
85 90 95

Ala Ser Lys Leu Leu Asp Phe Glu Ser Gln Arg
100 105

<210> 11

<211> 32

<212> DNA

<213> Artificial sequence

<220>

<223> primer PCR 1550

<400> 11

catgagatct ctataaaaac agagcggcta aa

32

<210> 12

<211> 42

<212> DNA

<213> Artificial sequence

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<223> primer PCR 1551

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tgacgcatgc actagtcata tgatgttcct tgttttttga tg

42

<210> 13

<211> 42

<212> DNA

<213> artificial sequence

<220>

<223> linker

<400> 13

tgacgcatgc actagtcata tgatgttcct tgttttttga tg

42

<210> 14

<211> 22

<212> DNA

<213> artificial sequence

<220>

<223> primer PCR 2386

<400> 14

gctcgggtacc cgggtgacta ac

22

<210> 15

<211> 27

<212> DNA

<213> artificial sequence

<220>

<223> primer PCR 2387

<400> 15

cttcccccg gcattattcc ctccagg

27

<210> 16
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 <212> DNA
 <213> artificial sequence

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 <223> primer PCR 2388
 <400> 16
 ccatcgatct cacacgctta gacgctaa 28

 <210> 17
 <211> 29
 <212> DNA
 <213> artificial sequence

 <220>
 <223> primer PCR 2389
 <400> 17
 ggactagtct aagttaaaag ccttaagat 29

 <210> 18
 <211> 26
 <212> DNA
 <213> artificial sequence

 <220>
 <223> primer PCR 2391
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 cgcggatcct tttaagaaag gtgttt 26

 <210> 19
 <211> 27
 <212> DNA

<213> artificial sequence

<220>

<223> primer PCR 2392

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<210> 20

<211> 32

<212> DNA

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<223> primer PCR 1777

<400> 20
gggaattcca tatgaatatac aaattaaagg at

32

<210> 21

<211> 35

<212> DNA

<213> Artificial sequence

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<223> primer PCR 1669

<400> 21
atcgcgatc cctagcttat cccaataaa tcctt

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<210> 22

<211> 34

<212> DNA

<213> Artificial Sequence

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<223> primer PCR 1783

<400> 22
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<210> 23

<211> 32

<212> DNA

<213> artificial sequence

<220>

<223> primer PCR 1784

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<210> 24

<211> 37

<212> DNA

<213> artificial sequence

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<223> primer PCR 1585

<400> 24
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<210> 25

<211> 35

<212> DNA

<213> artificial sequence

<220>

<223> primer PCR 1586

<400> 25
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<210> 26

<211> 20
<212> DNA
<213> artificial sequence

<220>
<223> oligonucleotide 2564
<400> 26
aatgtcgttt cggcttctga

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<210> 27
<211> 18
<212> DNA
<213> artificial sequence

<220>
<223> oligonucleotide 2565
<400> 27
taaaagcctt aagatatt

18

<210> 28
<211> 18
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<220>
<223> oligonucleotide H276f
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18

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<223> oligonucleotide H676r

<400> 29

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19